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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
09/380,447	09/01/1999	Sachdev S. Sidhu	11669.141USWO	2633
23552 MERCHANT 6	7590 08/30/2007 & GOULD PC		EXAM	INER
P.O. BOX 2903	3		STEELE, AMBER D	
MINNEAPOLIS, MN 55402-0903		*RED	ART UNIT	PAPER NUMBER
		KWK	1639	
			MAIL DATE	DELIVERY MODE
			08/30/2007	PAPER

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

Seq lesp 1: 09/30/2007 Seq lesp 3/PTA: 11/30/2007 Seq lesp stix: 02/29/2008



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APPLICATION NO. 09380447

FILING DATE 9/1/1999

FIRST NAMED INVENTOR SIDHU ET AL.

ATTORNEY DOCKET NO. 11669.141USWO

MERCHANT & GOULD PC P.O. BOX 2903 MINNEAPOLIS, MN 55402-0903

EXAMINER Amber D., Steele **ART UNIT PAPER**

20070821

DATE MAILED:

1639

Please find below and/or attached an Office communication concerning this application or proceeding.

Commissioner for Patents

- 1)The present application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 CFR 1.821 (a)(1) and (a)(2). However, this application fails to comply with the requirements of 37 CFR 1.821 through 1.825. The sequence listing has various errors as indicated in the attached Raw Sequence Listing and error report.
- 2)Applicant is given ONE MONTH from the date of this letter within which to comply with the Sequence Rules, 37 CFR 1.821-1.825. Failure to comply with these requirements will result in ABANDONMENT of the application under 37 CFR 1.821(g). Extensions of time may be obtained by filing a petition accompanied by the extension fee under the provisions of 37 CFR 1.136. In NO case may an applicant extend the period for response beyond the six month statutory period. Applicant is requested to return a copy of the attached Notice to Comply with the response.
- 3) Any inquiry concerning this communication or earlier communications from the examiner should be directed to Amber D. Steele whose telephone number is 571-272-5538. The examiner can normally be reached Monday through Friday 9:00AM-5:00PM. If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Doug Schultz can be reached at 571-272-0763. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300. Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

ADS August 21, 2007 Muhhally MARK L. SHIBUYA PRIMARY EXAMINER

PTO-90C (Rev.04-03)

Application No. Applicant(s) IDHU ET AL. 09/380,447 **Notice to Comply** Art Unit Examiner Steele, A. D. 1639

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE **DISCLOSURES**

Applicant must file the items indicated below within the time period set in the Office action to which the Notice is attached to avoid abandonment under 35 U.S.C. § 133 (extensions of time may be obtained under the provisions of 37 CFR 1 136(a))

the provisions of or it is too (e)).		
The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):		
I. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to the final rulemaking notice published at 55 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1990). If the effective filing date is on or after July 1, 1998, see the final rulemaking notice published at 63 FR 29620 (June 1, 1998) and 1211 OG 82 (June 23, 1998).		
 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c). 		
3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).		
5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).		
☐ 6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).		
□ 7. Other: see attached PTO-90C.		
Applicant Must Provide: ☑ An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".		
An initial or substitute paper copy of the "Sequence Listing", as well as an amendment specifically directing its entry into the application.		
A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).		
For questions regarding compliance to these requirements, please contact:		
For Rules Interpretation, call (571) 272-2510 For CRF Submission Help, call (571) 272-2501/2583. PatentIn Software Program Support		
Technical Assistance703-287-0200 To Purchase Patentin Software703-306-2600		
PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR REPLY		

Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Mon Jul 30 14:06:26 EDT 2007

Reviewer Comments:

<150> US 60/134,870

<151> 1999 05 19

<150> US 60/133,296

<151> 1999 05 10

<150> US 60/103,514

<151> 1998 10 08

<150> US 60/094,291

<151> 1998 07 27

<150> PCT/USUS99/16596

<151> 1999 07 22

The above non-ASCII characters ("squares") between dates appear throughout the submitted sequence listing file; they also appear in <222> responses which indicate locations within the sequence. Please replace them with hyphens.

<400> 293

His His His His

1

69

Please remove the above "69" appearing at the end of the submitted file.

Validated By CRFValidator v 1.0.2

Application No:

09380447

Version No:

3.0

Input Set:

Output Set:

Started: 2007-07-19 18:29:54.051

Finished: 2007-07-19 18:31:26.799

Elapsed:

0 hr(s) 1 min(s) 32 sec(s) 748 ms

Total Warnings: 284

Total Errors:

366

No. of SeqIDs Defined:

292

Actual SeqID Count:

293

Erro	r code	Error Description
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E	287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD)in <151>
E	287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD)in <151>
E	287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD)in <151>
E	287	Invalid WIPO ST.2 date format; Use (YYYY-MM-DD)in <151>
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W	213	Artificial or Unknown found in <213> in SEQ ID (1)
E	341	'Xaa' position not defined SEQID (1) POS (12)
E	341	'Xaa' position not defined SEQID (1) POS (13)
E	341	'Xaa' position not defined SEQID (1) POS (14)
E	341	'Xaa' position not defined SEQID (1) POS (15)
E	341	'Xaa' position not defined SEQID (1) POS (16)
E	341	'Xaa' position not defined SEQID (1) POS (17)
E	341	'Xaa' position not defined SEQID (1) POS (18)
E	341	'Xaa' position not defined SEQID (1) POS (19)
E	341	'Xaa' position not defined SEQID (1) POS (20)
E	341	'Xaa' position not defined SEQID (1) POS (21)
E	341	'Xaa' position not defined SEQID (1) POS (22)
E	341	'Xaa' position not defined SEQID (1) POS (23)
E	341	'Xaa' position not defined SEQID (1) POS (24)

Input Set:

Output Set:

Started: 2007-07-19 18:29:54.051

Finished: 2007-07-19 18:31:26.799

Elapsed: 0 hr(s) 1 min(s) 32 sec(s) 748 ms

Total Warnings: 284

Total Errors: 366

No. of SeqIDs Defined: 292

Actual SeqID Count: 293

Err	or code	Error Description
Έ	341	'Xaa' position not defined SEQID (1) POS (25)
E	341	'Xaa' position not defined SEQID (1) POS (26)
E	341	'Xaa' position not defined SEQID (1) POS (27)
E	341	'Xaa' position not defined SEQID (1) POS (28)
E	341	'Xaa' position not defined SEQID (1) POS (29)
E	341	'Xaa' position not defined SEQID (1) POS (30)
E	257	Invalid sequence data feature in <221> in SEQ ID (2)
E	257	Invalid sequence data feature in <221> in SEQ ID (3)
E	257	Invalid sequence data feature in $\langle 221 \rangle$ in SEQ ID (4)
E	257	Invalid sequence data feature in <221> in SEQ ID (5)
E	257	Invalid sequence data feature in <221> in SEQ ID (6)
E	257	Invalid sequence data feature in <221> in SEQ ID (7)
E	257	Invalid sequence data feature in <221> in SEQ ID (8)
W	213	Artificial or Unknown found in <213> in SEQ ID (9)
W	213	Artificial or Unknown found in <213> in SEQ ID (10)
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W	213	Artificial or Unknown found in <213> in SEQ ID (12)
W	213	Artificial or Unknown found in <213> in SEQ ID (13)
W	213	Artificial or Unknown found in <213> in SEQ ID (14)
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W	213	Artificial or Unknown found in <213> in SEQ ID (16)
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Input Set:

Output Set:

Started: 2007-07-19 18:29:54.051

Finished: 2007-07-19 18:31:26.799

Elapsed: 0 hr(s) 1 min(s) 32 sec(s) 748 ms

Total Warnings: 284
Total Errors: 366

No. of SeqIDs Defined: 292

Actual SeqID Count: 293

Error code	Error Description
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W 213	Artificial or Unknown found in <213> in SEQ ID (20)
W 213	Artificial or Unknown found in <213> in SEQ ID (21)
E 342	'n' position not defined found at POS: 20 SEQID(21)
E 342	'n' position not defined found at POS: 22 SEQID(21)
E 342	'n' position not defined found at POS: 26 SEQID(21)
E 342	'n' position not defined found at POS: 28 SEQID(21)
E 342	'n' position not defined found at POS: 31 SEQID(21)
E 342	'n' position not defined found at POS: 34 SEQID(21)
E 342	'n' position not defined found at POS: 38 SEQID(21)
Е 342	'n' position not defined found at POS: 41 SEQID(21)
E 342	'n' position not defined found at POS: 44 SEQID(21)
E 342	'n' position not defined found at POS: 47 SEQID(21)
W 213	Artificial or Unknown found in <213> in SEQ ID (22)
E 342	'n' position not defined found at POS: 19 SEQID(22)
.E 342	'n' position not defined found at POS: 22 SEQID(22)
E 342	'n' position not defined found at POS: 26 SEQID(22)
E 342	'n' position not defined found at POS: 28 SEQID(22)
E 342	'n' position not defined found at POS: 31 SEQID(22)
E 342	'n' position not defined found at POS: 35 SEQID(22)
E 342	'n' position not defined found at POS: 38 SEQID(22)

Input Set:

Output Set:

Started: 2007-07-19 18:29:54.051

Finished: 2007-07-19 18:31:26.799

Elapsed: 0 hr(s) 1 min(s) 32 sec(s) 748 ms

Total Warnings: 284

Total Errors: 366

No. of SeqIDs Defined: 292

Actual SeqID Count: 293

Erro	or code	Error Description
Ε	342	'n' position not defined found at POS: 41 SEQID(22)
E	342	'n' position not defined found at POS: 44 SEQID(22)
E	342	'n' position not defined found at POS: 46 SEQID(22) This error has occured more than 20 times, will not be displayed
W	213	Artificial or Unknown found in <213> in SEQ ID (23)
W	213	Artificial or Unknown found in <213> in SEQ ID (24)
W	213	Artificial or Unknown found in <213> in SEQ ID (25)
W	213	Artificial or Unknown found in <213> in SEQ ID (26)
W	213	Artificial or Unknown found in $<213>$ in SEQ ID (27) This error has occured more than 20 times, will not be displayed
E	257	Invalid sequence data feature in <221> in SEQ ID (36)
E	257	Invalid sequence data feature in <221> in SEQ ID (56)
E	355	Empty lines found between the amino acid numbering and the
E	321	No. of Bases conflict, this line has no nucleotides SEQID (293)
E	252	Calc# of Seq. differs from actual; 292 seqIds defined; count=293

Sequence Listing

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<110> Sidhu, Sachdev S.
     Weiss, Gregory A.
     Wells, James A.
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     COAT PROTEIN
<130> 11669.141USWO
<140> 09380447
<141> 1999-09-01
<150> US 09/380,447
<151> 1999 09 01
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<150> US 60/133,296
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<151> 1999 07 22
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<222> 12 30
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40

35

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                 20
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                20
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<213> Zj 2 phage
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                       10 15
 1 5
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               20
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<222> 1 50

<223> coat protein VIII

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<221> Ike phage
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                                    10
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                  20
                                     25
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                                     40
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 atcgtc 56
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 aggtgtcgtg g 61
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 taactccctg caagcc 66
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                                      25
 Tyr Met Leu Leu Val Glu Ala Ser Pro Trp Ala Ala Lys Ala Pro
                  35
                                      40
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